



4/25

A

ZNFN3A1 : 1 MEPLKVEKFATANRGNGLRAVTPLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM 60
ME LKVEKF TANRGNGLRAV PLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM
AK010447: 1 MEALKVEKFTTANRGNGLRAVAPLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM 60

ZNFN3A1 : 61 RCSQCRVAKYCSAKCQKKAWPDHKRECKLSCKPRYPDSVRL LGRVVKLMDGAPSES 120
RCSQCR+AKYCSAKCQKKAWPDH+REC CLKCKPRYPDSVRL LGRV+ KLMD PSES B
AK010447: 61 RCSQCRVAKYCSAKCQKKAWPDHRECSCLCKPRYPDSVRL LGRVIVKLMDEKPSES 120

ZNFN3A1 : 121 EKLYSFYDLESNINKLTEDKKEGLROLMTFOHFMRREEIQDASQLPPAFDLFEAFKVIC 180
EKLYSFYDLESNI+KLTEDKKEGLROLMTFOHFMRREEIQDASQLPP+FDLFEAFKVIC
AK010447: 121 EKLYSFYDLESNISKLTEDKKEGLROLAMTFOHFMRREEIQDASQLPPSFDLFEAFKVIC 180

ZNFN3A1 : 181 NSFTICNAEMQEVGVGLYPSISLNHSCDPNCSIVFNGPHLLRAVRDIEVGEEITICYL 240
NSFTICNAEMQEVGVGLYPS+SLNHSCDPNCSIVFNGPHLLRAVR+IE GEEITICYL
AK010447: 181 NSFTICNAEMQEVGVGLYPSMSLNHSCDPNCSIVFNGPHLLRAYREIEAGEEITICYL 240

ZNFN3A1 : 241 DMLMTSEERRKQLRDQYCFE CDCFRCQTQDKDADMLTGDEQVWKEVQESLKKIEELKAHW 300
DMLMTSEERRKQLRDQYCFE CDC RCQTQDKDADMLTGDEQ+WKEVQESLKKIEELKAHW
AK010447: 241 DMLMTSEERRKQLRDQYCFE CDCIRCQTQDKDADMLTGDEQIWKEVQESLKKIEELKAHW 300

ZNFN3A1 : 301 KWEQVLAMCQAISSNSERLPDINIYQLKVLDACINLGLLEEALFYGTRTMEPYRI 360
KWEQVLA+CQAI+SNS RLPDINIYQLKVLDACINLG+LEEALFY RTMEPYRI
AK010447: 301 KWEQVLALCQAIINSNSNRLPDINIYQLKVLDACINLGMLEEALFYAMRTMEPYRI 360

ZNFN3A1 : 361 FFPGSHPVGRGVQVMKVGKLQLHQGMFPQAMKNLRLAFDIMRVTHGREHSLIEDLILLEE 420
FFPGSHPVGRGVQVMKVGKLQLHQGMFPQAMKNLRLAFDIM+VTHGREHSLIEDLILLEE
AK010447: 361 FFPGSHPVGRGVQVMKVGKLQLHQGMFPQAMKNLRLAFDIMKVTHGREHSLIEDLILLEE 420

ZNFN3A1 : 421 CDANIRAS 428 (SEQ ID NO: 1)
CDANIRAS (SEQ ID NO: 81)
AK010447: 421 CDANIRAS 428 (SEQ ID NO: 82)

FIG.2B

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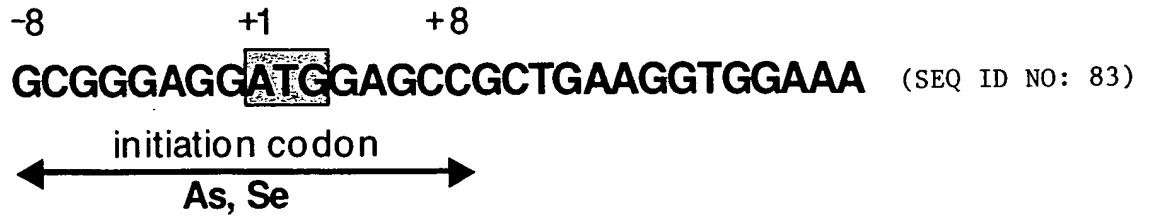


FIG.6A

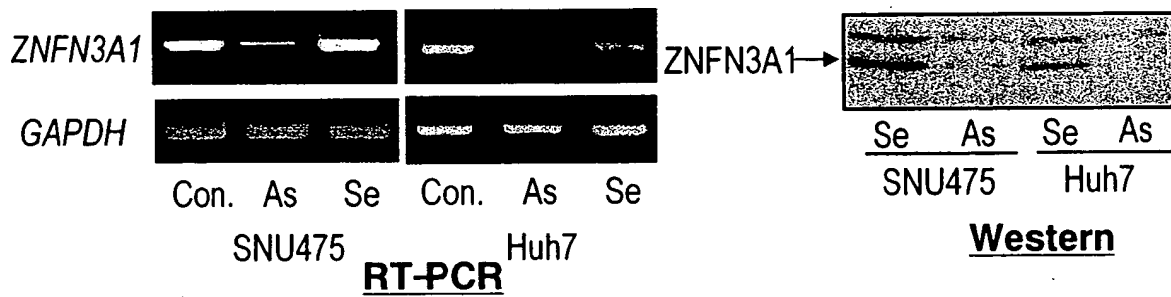


FIG.6B

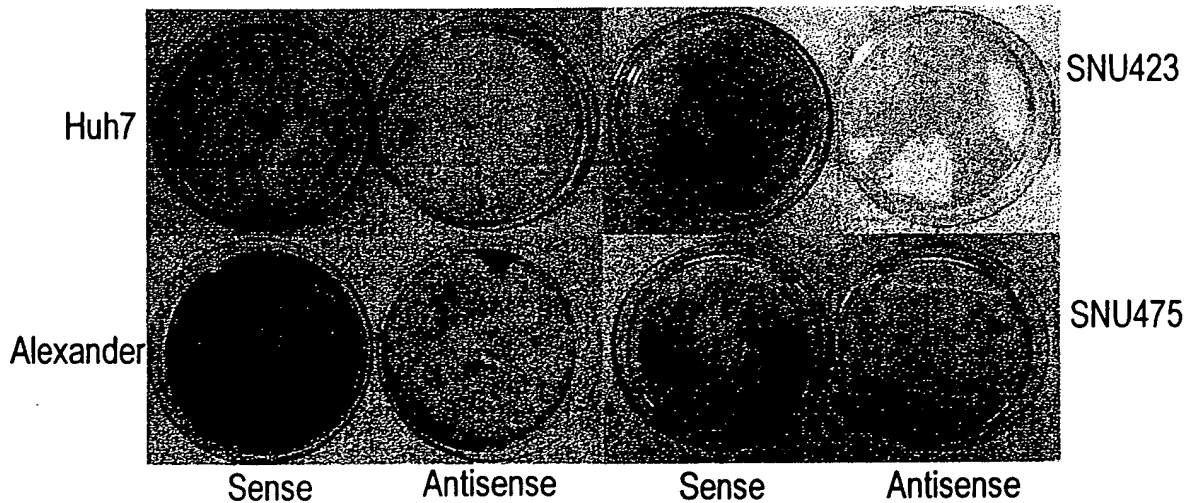


FIG.6C